



RAW SEQUENCE LISTING ERROR REPORT

The Biotechnology Systems Branch of the Scientific and Technical Information Center (STIC) detected errors when processing the following computer readable form:

Application Serial Number: 10/050,000 A

Source: PCT

Date Processed by STIC: 11/5/03

THE ATTACHED PRINTOUT EXPLAINS DETECTED ERRORS.

PLEASE FORWARD THIS INFORMATION TO THE APPLICANT BY EITHER:

- 1) INCLUDING A COPY OF THIS PRINTOUT IN YOUR NEXT COMMUNICATION TO THE APPLICANT, WITH A NOTICE TO COMPLY or,
- 2) TELEPHONING APPLICANT AND FAXING A COPY OF THIS PRINTOUT, WITH A NOTICE TO COMPLY

FOR CRF SUBMISSION AND PATENTIN SOFTWARE QUESTIONS, PLEASE CONTACT MARK SPENCER, TELEPHONE: 703-308-4212; FAX: 703-308-4221

Effective 12/13/03: TELEPHONE: 571-272-2510; FAX: 571-273-0221

TO REDUCE ERRORED SEQUENCE LISTINGS, PLEASE USE THE CHECKER VERSION 4.1 PROGRAM, ACCESSIBLE THROUGH THE U.S. PATENT AND TRADEMARK OFFICE WEBSITE. SEE BELOW FOR ADDRESS:

<http://www.uspto.gov/web/offices/pac/checker/chkr41note.htm>

Applicants submitting genetic sequence information electronically on diskette or CD-Rom should be aware that there is a possibility that the disk/CD-Rom may have been affected by treatment given to all incoming mail.

Please consider using alternate methods of submission for the disk/CD-Rom or replacement disk/CD-Rom.

Any reply including a sequence listing in electronic form should NOT be sent to the 20231 zip code address for the United States Patent and Trademark Office, and instead should be sent via the following to the indicated addresses:

1. EFS-Bio (<http://www.uspto.gov/ebs/efs/downloads/documents.htm>), EFS Submission User Manual - ePAVE)
2. U.S. Postal Service: Commissioner for Patents, P.O. Box 1450, Alexandria, VA 22313-1450
3. Hand Carry directly to (EFFECTIVE 12/01/03):
U.S. Patent and Trademark Office, Box Sequence, Customer Window, Lobby, Room 1B03, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202
4. Federal Express, United Parcel Service, or other delivery service to: U.S. Patent and Trademark Office, Box Sequence, Room 1B03-Mailroom, Crystal Plaza Two, 2011 South Clark Place, Arlington, VA 22202

Revised 10/08/03

Raw Sequence Listing Error Summary

<u>ERROR DETECTED</u>	<u>SUGGESTED CORRECTION</u>	SERIAL NUMBER: <u>10/050,000 A</u>
ATTN: NEW RULES CASES: PLEASE DISREGARD ENGLISH "ALPHA" HEADERS, WHICH WERE INSERTED BY PTO SOFTWARE		
1 <u> </u> Wrapped Nucleics Wrapped Aminos	The number/text at the end of each line "wrapped" down to the next line. This may occur if your file was retrieved in a word processor after creating it. Please adjust your right margin to .3; this will prevent "wrapping."	
2 <u> </u> Invalid Line Length	The rules require that a line not exceed 72 characters in length. This includes white spaces.	
3 <u> </u> Misaligned Amino Numbering	The numbering under each 5 th amino acid is misaligned. Do not use tab codes between numbers; use space characters, instead.	
4 <u> </u> Non-ASCII	The submitted file was not saved in ASCII(DOS) text, as required by the Sequence Rules. Please ensure your subsequent submission is saved in ASCII text.	
5 <u> </u> Variable Length	Sequence(s) <u> </u> contain n's or Xaa's representing more than one residue. Per Sequence Rules, each n or Xaa can only represent a single residue. Please present the maximum number of each residue having variable length and indicate in the <220>-<223> section that some may be missing.	
6 <u> </u> PatentIn 2.0 "bug"	A "bug" in PatentIn version 2.0 has caused the <220>-<223> section to be missing from amino acid sequences(s) <u> </u> . Normally, PatentIn would automatically generate this section from the previously coded nucleic acid sequence. Please manually copy the relevant <220>-<223> section to the subsequent amino acid sequence. This applies to the mandatory <220>-<223> sections for Artificial or Unknown sequences.	
7 <u> </u> Skipped Sequences (OLD RULES)	Sequence(s) <u> </u> missing. If intentional, please insert the following lines for each skipped sequence: (2) INFORMATION FOR SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) (i) SEQUENCE CHARACTERISTICS: (Do not insert any subheadings under this heading) (xi) SEQUENCE DESCRIPTION:SEQ ID NO:X: (insert SEQ ID NO where "X" is shown) This sequence is intentionally skipped Please also adjust the "(ii) NUMBER OF SEQUENCES:" response to include the skipped sequences.	
8 <u> </u> Skipped Sequences (NEW RULES)	Sequence(s) <u> </u> missing. If intentional, please insert the following lines for each skipped sequence. <210> sequence id number <400> sequence id number 000	
9 <u> </u> Use of n's or Xaa's (NEW RULES)	Use of n's and/or Xaa's have been detected in the Sequence Listing. Per 1.823 of Sequence Rules, use of <220>-<223> is MANDATORY if n's or Xaa's are present. In <220> to <223> section, please explain location of n or Xaa, and which residue n or Xaa represents.	
10 <u> </u> Invalid <213> Response	Per 1.823 of Sequence Rules, the only valid <213> responses are: Unknown, Artificial Sequence, or scientific name (Genus/species). <220>-<223> section is required when <213> response is Unknown or is Artificial Sequence	
11 <u> </u> Use of <220>	Sequence(s) <u> </u> missing the <220> "Feature" and associated numeric identifiers and responses. Use of <220> to <223> is MANDATORY if <213> "Organism" response is "Artificial Sequence" or "Unknown." Please explain source of genetic material in <220> to <223> section. (See "Federal Register," 00/01/1998, Vol. 63, No. 104, pp. 29631-32) (Sec. 1.823 of Sequence Rules)	
12 <u> </u> PatentIn 2.0 "bug"	Please do not use "Copy to Disk" function of PatentIn version 2.0. This causes a corrupted file, resulting in missing mandatory numeric identifiers and responses (as indicated on raw sequence listing). Instead, please use "File Manager" or any other manual means to copy file to floppy disk.	
13 <u> </u> Misuse of n/Xaa	"n" can only represent a single <u>nucleotide</u> ; "Xaa" can only represent a single <u>amino acid</u>	



PCT

RAW SEQUENCE LISTING

DATE: 01/05/2004

PATENT APPLICATION: US/10/050,000A

TIME: 11:45:56

Input Set : A:\2002-07-07 0365-0529P.ST25.txt

Output Set: N:\CRF4\01052004\J050000A.raw

3 <110> APPLICANT: NAKARI-SETALA, Tiina
 5 <120> TITLE OF INVENTION: A METHOD FOR DECREASING THE FOAM FORMATION DURING
 CULTIVATION OF A
 6 MICROORGANISM
 8 <130> FILE REFERENCE: 0365-0529P
 C--> 10 <140> CURRENT APPLICATION NUMBER: US/10/050,000A
 C--> 11 <141> CURRENT FILING DATE: 2002-07-10
 13 <160> NUMBER OF SEQ ID NOS: 8
 15 <170> SOFTWARE: PatentIn Ver. 2.1
 17 <210> SEQ ID NO: 1
 18 <211> LENGTH: 2868
 19 <212> TYPE: DNA
 20 <213> ORGANISM: Trichoderma reesei
 22 <220> FEATURE:
 23 <221> NAME/KEY: gene
 24 <222> LOCATION: (1523)..(1950)
 25 <223> OTHER INFORMATION: hfb1
 27 <400> SEQUENCE: 1
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 30 cgtatcccat tgctgggctg aaagcttcac acgtatcgca taagcgtctc caaccagtgc 180
 31 ttagggtgacc cttacaggata cttacagtaa gactgtatta agtcagtccac tctttcactc 240
 32 gggctttgaa tacgatacctc aatactcccg ataacagtaa gaggatgata cagcctgcag 300
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 36 cccgttaaac tcccctgtaa cttggcatca ctcatctgtg atcccaacag actgagttgg 540
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 39 cccggacgca ccacaactcc cctgtgtca ttgagtcccc atatgcgtct tctcagcgtg 720
 40 caactctgag acggattagt cctcacgatg aaattaactt ccagcttaag ttcgtagcct 780
 41 tgaatgagtg aagaaatttc aaaaacaaac tgagtagagg tcttgagcag ctgggggtgg 840
 42 acgcccctcc tgcactcttg ggacatcgta cggcagagaa tcaacggatt cacacctttg 900
 43 ggtcgagatg agctgatctc gacagatacg tgcttcacca cagctgcagc tacctttgcc 960
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 49 ttgagattgt tgaccgggaa aggggggacg aagaggatgg ctgggtgagg tggattggg 1320
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 51 ctgtcttctc aaaatctcct tccatcttgt ccttcacag caccagagcc agcctgaaca 1440
 52 cctccagtca acttccctta ccagtacatc tgaatcaaca tccattcttt gaaatctcac 1500
 53 cacaaccacc atcttcttca aaatgaagtt cttcgccatc gccgtctctt ttgccgccgc 1560

**Does Not Comply
 Corrected Diskette Needed**

(pg. 3)

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DATE: 01/05/2004

PATENT APPLICATION: US/10/050,000A

TIME: 11:45:56

Input Set : A:\2002-07-07 0365-0529P.ST25.txt

Output Set: N:\CRF4\01052004\J050000A.raw

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58 gcgcaaaaac cggcgcccag cctctctgct gcgtggcccc cggttgtaagt tgatgcccc 1860
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63 caatatttca gtacactttt cttcataaat caaaaagact gctattctct ttgtgacatg 2160
64 ccggaaggga acaattgctc ttggtctctg ttatttgcaa gtaggagtgg gagattcgcc 2220
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68 ttcagctgat atccacgcca atgatacagc gatatacctc caatatctgt gggaacgaga 2460
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72 aacgttacta caaaagacat gggcgaatgc aaagacctga aagcaaacc tttttgcgac 2700
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74 tgagataata catgaaaagc acaacggaaa cgaaagaacc aggaaaagaa taaatctatc 2820
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79 <211> LENGTH: 3585
80 <212> TYPE: DNA
81 <213> ORGANISM: Trichoderma reesei
83 <220> FEATURE:
84 <221> NAME/KEY: gene
85 <222> LOCATION: (1191)..(1593)
86 <223> OTHER INFORMATION: hfb2
88 <400> SEQUENCE: 2
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91 cgctttgtat tattagagta tatgcaagtc tcaggactat cgactcaact ctaccacccg 180
92 aggacgatcg cggcacgata cgccctcggt ctcatgtgcc caagcagacc aactgccct 240
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94 cactccctt tggccaaatc agcttgcatg tcaagagaca tcgagctgtg cttgaaatt 360
95 actaacaacc agggatggga aacgaagcct gcttttgga agacaacaat gagagagaga 420
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97 aatgtcacat ccgagtcttg gggcctctgt gagaatgtcc agagtaatac gtgttttgcg 540
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99 tttcgaagta ttatctggag gatagaagac gtctaggtaa ctacacaaa ggctataact 660
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101 cagcctcttg gagcgtcat agatcacatc tgtgtagacc gacattctat gaataatcat 780
102 ctcatcatgg ccacatacta ctacatacgt gtctctgcct acctgacatg tagcagtggc 840
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104 gagagattgc gatgagccct ctccctacct acagacggct gacaatgtcc gtataccacc 960
105 agccaacgtg atgaaaacaa ggacatgagg aacagcctgc gagagctgga agatgaagag 1020
106 ggccagaaaa aaaagtataa agaagacctc gattcccgcc atccaacaat cttttccatc 1080

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DATE: 01/05/2004

PATENT APPLICATION: US/10/050,000A

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Input Set : A:\2002-07-07 0365-0529P.ST25.txt

Output Set: N:\CRF4\01052004\J050000A.raw

W--> 107 ctcatcagca cactcatcta caaccatcac cacattcact caactcctct ttctcaactc 1140
 108 tccaaacaca aacattcttt gttgaatacc aaccatcacc acctttcaag atgcagttct 1200
 109 tcgccgtcgc cctcttcgcc accagcgccc tggctgctgt ctgccctacc ggcctcttct 1260
 110 ccaaccctct gtgctgtgcc accaacgtcc tcgacctcat tggcggtgac tgcaagaccc 1320
 111 gtatgttgaa ttccaatctc tgggcatcct gacattggac gatacagttg acttacacga 1380
 112 tgctttacag ctaccatcgc cgtcgacact ggcgccatct tccaggctca ctgtgccagc 1440
 113 aagggtcca agcctctttg ctgctgtgct cccgtggtaa gtagtgctcg caatggcaaa 1500
 114 gaagtaaaaa gacatttggt cctgggatcg ctaactcttg atatcaaggc cgaccaggct 1560
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 118 tttgcgagca ggacatagac gtatctagag ttctagtcaa tacattattg aaaagttgga 1800
 119 gtatacctat cgctggcact ggtatcttga agatatcttc tcttctgtg aggttatgta 1860
 120 **tggaatcag tcgaaatcta tttgaagaca gagctcaage ttcaaacatt cacctgga** 1920
 121 ttgaccattt tgtttcgatg gttgcagttg gtgggtgtca cttctgcaat catgtacgag 1980
 122 cacaagtata gcagtattcc atctgatctg catctgggta aatgtcgcca ctctacctag 2040
 123 gtacccaata aataccgaat tggtcagctc tcgggtgaca aaccggcccg cttttcgacc 2100
 124 gtgctctgtc caattctagg ctgttcaatg gttctgact gtgataaacc ttggagcta 2160
 125 cataacttac cttacaataa atccaactgc cggcacttgc ttcccttcac ccaaccactc 2220
 126 gcaaacatca cgcaacctgt ctcgatcccc tgtccgaaat ctgcttgga acgtatcatc 2280
 127 acaaatcata cacacagaca aaaaggagcc aaagcagcaa tggcaagaca ccgaggccgg 2340
 128 cagcgcgccc gtcgccgttt ttaaaaagcg aagcgcaaa ggcaaagcca acctgcgcaa 2400
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 130 cgtcggaaga cgaagccggg cacagagtca agaggcgcaa gaggacggc gtgctaccg 2520
 131 ccgccgcca ggggcgcgc gccagcaac cgggacgacg gcggcgccg aacagccgc 2580
 132 ttcacggcca acagaagcgt cccgattgct gacagcaacg acgcgacca gcacagcaac 2640
 133 tggtagcagc aggacgcaaa ggacgcgctc tcggcaaaga acctcctcg atcttcgaga 2700
 134 gcgtccaagg acgcgcagcc agacggcagc tacaagggcc tggcgaacca gacgtcctt 2760
 135 atacaaaaga atccggatgc gcccggaag acagttggc ccgtcaagg gcctaccaac 2820
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 140 cgcgagagaa gacctcaagg caggctggca gctggatcaa gaggggaaa aggtcaccaa 3120
 141 gggcaagaag aacctggggg gaacggtagt ggccagcgcg aaccggaaca aggccaaggt 3180
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 146 gagggcgccc aggaggagac aggaggcgat agagaggggc gaggaagtca gtgatgaaga 3480
 147 ggaggaggag gaggaggact gatgatgatg gggcagatg acgatgcagg tcgactctag 3540
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 151 <210> SEQ ID NO: 3
 152 <211> LENGTH: 63
 153 <212> TYPE: DNA
 154 <213> ORGANISM: Artificial Sequence
 156 <220> FEATURE:
 157 <223> OTHER INFORMATION: Description of Artificial Sequence: 5' primer
 159 <400> SEQUENCE: 3

- Please explain "N" locations. Please see item #9 on error summary sheet and explanation on page 5.

RAW SEQUENCE LISTING

DATE: 01/05/2004

PATENT APPLICATION: US/10/050,000A

TIME: 11:45:56

Input Set : A:\2002-07-07 0365-0529P.ST25.txt

Output Set: N:\CRF4\01052004\J050000A.raw

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161 ggc 63
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165 <211> LENGTH: 30
166 <212> TYPE: DNA
167 <213> ORGANISM: Artificial Sequence
169 <220> FEATURE:
170 <223> OTHER INFORMATION: Description of Artificial Sequence: 3'primer
172 <400> SEQUENCE: 4
173 tcgtacggat cctcaagcac cgacggcgg 30
176 <210> SEQ ID NO: 5
177 <211> LENGTH: 13
178 <212> TYPE: PRT
179 <213> ORGANISM: Artificial Sequence
181 <220> FEATURE:
182 <223> OTHER INFORMATION: Description of Artificial Sequence: linker
184 <400> SEQUENCE: 5
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186 1 5 10
189 <210> SEQ ID NO: 6
190 <211> LENGTH: 14
191 <212> TYPE: PRT
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194 <220> FEATURE:
195 <223> OTHER INFORMATION: Description of Artificial Sequence: linker
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199 1 5 10
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205 <213> ORGANISM: Artificial Sequence
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208 <223> OTHER INFORMATION: Description of Artificial Sequence: 5'primer
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215 <211> LENGTH: 34
216 <212> TYPE: DNA
217 <213> ORGANISM: Artificial Sequence
219 <220> FEATURE:
220 <223> OTHER INFORMATION: Description of Artificial Sequence: 3'primer
222 <400> SEQUENCE: 8
223 tgaattccat atgctaaccc cgtttcatct ccag 34

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RAW SEQUENCE LISTING ERROR SUMMARY
PATENT APPLICATION: US/10/050,000A

DATE: 01/05/2004
TIME: 11:45:57

Input Set : A:\2002-07-07 0365-0529P.ST25.txt
Output Set: N:\CRF4\01052004\J050000A.raw

Please Note:

Error Explanation:

Use of n and/or Xaa have been detected in the Sequence Listing. Please review the Sequence Listing to ensure that a corresponding explanation is presented in the <220> to <223> fields of each sequence which presents at least one n or Xaa.

Seq#:2; N Pos. 1917,2160,3515

VERIFICATION SUMMARY

PATENT APPLICATION: US/10/050,000A

DATE: 01/05/2004

TIME: 11:45:57

Input Set : A:\2002-07-07 0365-0529P.ST25.txt

Output Set: N:\CRF4\01052004\J050000A.raw

L:10 M:270 C: Current Application Number differs, Replaced Current Application Number
L:11 M:271 C: Current Filing Date differs, Replaced Current Filing Date
L:120 M:341 W: (46) "n" or "Xaa" used, for SEQ ID#:2 after pos.:1860
M:341 Repeated in SeqNo=2